

SEQUENCE LISTING

- <110> PINEL Anne-Marie HOCQUAUX Michel
- <120> NOVEL PEPTIDIC CONJUGATES FOR ALOPECIA PREVENTIVE AND CURATIVE TREATMENT
- <130> 3493-0157PUS1
- <140> PCT/FR2004/001882
- <141> 2004-07-16
- <150> FR 03/08797
- <151> 2003-07-18
- <160> 12
- <170> PatentIn version 3.2
- <210> 1
- <211> 11
- <212> PRT
- <213> Artificial sequence
- <220>
- <223> Synthetic Peptide
- <220>
- <221> misc feature
- <222> (1)..(3)
- <223> Xaa can be Glu-Gln-Arg, Arg-Lys, Arg-Lys-Asp sequences or Arg amino acid or a bond and up to 3 residues may be present or absent
- <220>
- <221> misc feature
- <222> (7)..(11)
- <223> Xaa can be Tyr-Val-Gln-Leu-Tyr-Amide, Leu-DOPA sequences, the amino acids Dopa amide or HomoPhe amide and up to 4 residues may be present or absent
- <400> 1

Xaa Xaa Xaa Lys Asp Val Xaa Xaa Xaa Xaa Xaa 1 5 10

- <210> 2
- <211> 11
- <212> PRT
- <213> Artificial sequence
- <220>
- <223> Synthetic Peptide

```
<220>
<221> misc_feature
<222> (1)..(3)
<223> Xaa can be Gly-Gln-Gln or Glu-Gln sequences and up to
      1 residue may be present or absent
<220>
<221> misc feature
<222> (7)..(11)
<223> Xaa can be Tyr-Val-Gln-Leu-Tyr-Amide, Leu-DOPA, Val-Tyr,
      Val-Tyr-amide sequences, or the amino acids Tyr, Tyr amide,
      Dopa amide or HomoPhe amide amide and up to 4 residues may
      be present or absent
<400> 2
Xaa Xaa Xaa Lys Asp Val Xaa Xaa Xaa Xaa
<210> 3
<211> 11
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic Peptide
<220>
<221> MOD RES
<222> (1)..(1)
<223> BLOCKED
<220>
<221> misc feature
<222> (1)..(3)
<223> Xaa can be Glu-Gln-Arg, Arg-Lys, Arg-Lys-Asp sequences
      or Arg amino acid or a bond and up to 3 residues may
      be present or absent
<220>
<221> misc_feature
<222> (7)..(11)
<223> Xaa can be Tyr-Val-Gln-Leu-Tyr-Amide, Leu-DOPA sequences,
      the amino acids Dopa amide or HomoPhe amide and up to 4
      residues may be present or absent
<400> 3
Xaa Xaa Xaa Lys Asp Val Xaa Xaa Xaa Xaa Xaa
                5
<210> 4
<211> 11
<212> PRT
```

```
<213> Artificial sequence
<220>
<223> Synthetic Peptide
<220>
<221> MOD RES
<222> (1)..(1)
<223> BLOCKED
<220>
<221> misc_feature
<222> (1)..(3)
<223> Xaa can be Gly-Gln-Gln or Glu-Gln sequences and up to
      1 residue may be present or absent
<220>
<221> misc_feature
<222> (7)..(11)
<223> Xaa can be Tyr-Val-Gln-Leu-Tyr-Amide, Leu-DOPA, Val-Tyr,
      Val-Tyr-amide sequences, or the amino acids Tyr, Tyr
      amide, Dopa amide or HomoPhe amide and up to 4 residues
      may be present or absent
<400> 4
Xaa Xaa Xaa Lys Asp Val Xaa Xaa Xaa Xaa Xaa
                5
<210> 5
<211> 5
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic Peptide
<220>
<221> MOD_RES
<222> (1)..(1)
<223> BLOCKED
<220>
<221> MOD RES
<222> (5)..(5)
<223> AMIDATION
<220>
<221> misc feature
<222> (5)..(5)
<223> Xaa = homophenylalanine amide.
<400> 5
```

```
Arg Lys Asp Val Xaa
<210> 6
<211> 4
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic Peptide
<220>
<221> MOD RES
<222> (1)..(1)
<223> BLOCKED
<220>
<221> MOD_RES
<222> (4)..(4)
<223> AMIDATION
<220>
<221> misc_feature
<222> (4)..(4)
<223> Xaa = dihydrophenylalanine amide.
<400> 6
Lys Asp Val Xaa
<210> 7
<211> 5
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic Peptide
<220>
<221> MOD_RES
<222> (1)..(1)
<223> BLOCKED
<220>
<221> MOD_RES
<222> (5)..(5)
<223> AMIDATION
<220>
<221> misc feature
<222> (5)..(5)
<223> Xaa = dihydrophenylalaline amide.
<400> 7
```

```
Arg Lys Asp Val Xaa
<210> 8
<211> 6
<212> PRT
<213> Artificial sequence
<223> Synthetic Peptide
<220>
<221> misc_feature
<222> (1)..(3)
<223> Xaa is Lys or MeLys and up to two residues may be present
      or absent
<400> 8
Xaa Xaa Xaa Gly His Lys
                5
<210> 9
<211> 6
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic Peptide
<220>
<221> misc feature
<222> (1)..(3)
<223> Xaa is Lys or MeLys and up to two residues may be present
      or absent
<220>
<221> MOD_RES
<222> (6)..(6)
<223> AMIDATION
<400> 9
Xaa Xaa Gly His Lys
<210> 10
<211> 6
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic Peptide
```

```
<220>
<221> MOD_RES
<222> (1)..(1)
<223> BLOCKED
<220>
<221> misc_feature
<222> (1)..(3)
<223> Xaa is Lys, MeLys, or a bond and up to three residues may be
      present or absent
<400> 10
Xaa Xaa Xaa Gly His Lys
                5
<210> 11
<211> 6
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic Peptide
<220>
<221> MOD RES
<222> (1)..(1)
<223> BLOCKED
<220>
<221> misc_feature
<222> (1)..(3)
<223> Xaa is Lys, MeLys, or a bond and up to three residues may be
      present or absent
<220>
<221> MOD_RES
<222> (6)..(6)
<223> AMIDATION
<400> 11
Xaa Xaa Gly His Lys
                5
1
<210> 12
<211> 5
<212> PRT
<213> Artificial Sequence
<220>
<223> Synthetic peptide
```